Figure 1, page 1

EcoR I

- 4102 -~~4152~~ AGGAATTCAT CCATTTAAAT CATACAATTT AATGGCTTTT AGTATATTCA
- 4052 -~~4102~~ CAGGTTGTGC ATCCATCACA ATCCATTTTA GAACAGTTTT ATTACTCCAA
<HNF-3/Fkh-1 SREBP>
B2 - 4002 -~~4052~~ AAATAAACCC TGCATTCCTT AGCCATCACC CCCCAACATC CTCCATCCTC
- 3952
NF-Y>
-~~4002~~ CTTCCAAGCC CTGGGCAACC ACCAATCTAC TTTCTGTCTC TATAAATTTG
- 3902 -~~3952~~ CCAATTCTGG ACATTTCATA TAAATGGAAG CAAACAACAT GTGAGACTTT
<NF-Y <IRF-2
- 3852 -~~3902~~ GTGACTGGCT GCTTTCACTT AGCATTCTAT TTTTAAGGCT CATTATGTTA
- 3802 -~~3852~~ CAGTACTTAG CAGTACTTCA TTCTTTTTTA TTCTCAAATG GTATTCCACT
- 3752 -~~3802~~ GTGTGGGTAT CCCATATCAT ATTATTAGAG ACAGGTTCTC ACTCTGTCAC
- 3702 -~~3752~~ CCAGGCTGGA GTGCAGTGGC ACAATCATAG CTCACTGTAA CCTCAAACTC
<SREBP
- 3652 -~~3702~~ CTGGGCTCAA GTGATCCTAC TACCTCAGCC TCCAGAGTAG CTAGGACTAC
- 3602
<IRF-1
- 3552 -~~3652~~ AGGCACACAC AGCCATACCT GGCTAATTTT TTTTTTTAAT TTTCATTTTA
- 3502 -~~3602~~ TGTATTCATT TTCTTTCTTT TTTGTTGTTG TTGTTTTGAG ATAGGGTCTC
-~~3552~~ ACTTTGTTAC CCAGGCTGGA GGGCAGTGGC ATGGTGACAG CTGAGCAGCC
- 3452
<SREBP
-~~3502~~ TTGACTTCCT GGGCTCAAGT GATCCTCCTG CCTCAGCCTC CCAAGTAGCT
- 3402 -~~3452~~ GGGACTACAA ACACGTGTCA CCATGCCTGG CTGATATTTT TTTTCTTGAA
- 3352 -~~3402~~ ACAGGGTATC ACTCTGTTGC CCAGGCTGGA GTACAGTGGC GTAATAATAG
c1
Pst I

- 3302 -~~3352~~ CTCACTGCAG CCTCCCCTCC TGGGCTCAAG CAATCCGCTG GCCTCAGCAT
- 3252 -~~3302~~ CCTGAGTAGC TGGGACTACA GGCTTGTGCC ACCAGGCCCA GCTAAGTTTT
- 3202 -~~3252~~ AAAAAATGAT TTTTGGTATA GAGGAGGTCT TGCTATGTTG CTCAGGCTGT
SREBP>
- 3152 -~~3202~~ ATTTTTATTG TTGAGACAAG GTCTCACTAT GTTGCCATGA TCCCCCCACC
<AP-1
- 3102 -~~3152~~ TCCACTTCCC AAAGTGCTCA TCTTATCTGT TCATTAGTCA GTTGACAGAC
- 3052
<RAR-α1
-~~3102~~ ATTTAGGTTG TTTCCACTTT TTGACCATTA TGAATAATAC TCCAGTGAAT
- 3002 -~~3052~~ ATTCATGTAT ACATTTGTGT GGGCATATGT TTTCATTTCT GTTGGGTTTA
- 2952 -~~3002~~ TATCTAGGAG TGGAATTGCT GGATCCCGGG TAATATTTTG ACAGGCAGAG
- 2902
C/EBP-β>
- 2852 -~~2952~~ TTCAGGGGAA GAAAAACTTG GGAAAATGAA GCATGTTTAG AAATCAGCAA
- 2802 -~~2902~~ GAGTGCAGGG GTTTTTCGGA GTTTTATTTT ATATTCTGTT GACAAATGTG
-~~2852~~ CAGTTTGATG AAGATACAAG TTATACTAAG TGAGAAGTGA GAATTAAGGC
- 2752 -~~2802~~ TGGAATAGGG CGTTCAGAGT AAAATCATGA AGCACTTTGA ATACCAAAAT
NF-1> <HNF3-β
- 2702 -~~2752~~ TAAGGAGCTT GGCTGTAAAC AAAATAATAA AAAATCACAA TTTTTTTTTT
- 2652 -2~~702~~ TTTTTTGAGA AAGAGTCTTG CTCTTTCACC CTGGCTGGAG GGCAGTGGTG
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- 2602 -~~2652~~ TGATCTCAGC TCACTGCAAC TTTCGCCTCC CGGGTTCAAG CAATTCTCCT

Figure 1, page 2

<NF-kB

-2552 ~~-2602~~ GCTTCAGCCT CCCAAGTAGC TGGGACTACA GGCACTTCCC ACCATGCCCA

NF-kB><IRF-1

-2502 ~~-2552~~ GCTGATTTTT GTATTTTTAG TAGAGATGGG ATTTCACTTT GTTGGCCAAG

-2452 ~~-2502~~ CTGGTCTCAA ACTTTTTGCT GTCATAATTG TTGTAACTAT TGTTCCTTTT

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-2402 ~~-2452~~ GCTGAGGTAG GGCCCCCAGA CCAAAAAAAA TAAATCTTAG AATCCAAATC

-2352 ~~-2402~~ AGTGTGTTGG TTTGACCACT GTCACTTGAG AACCACAGTG TGACCAGGGC

B2

C2

Taq I

---- IRF-2>

-2302 ~~-2352~~ CTCAGGAGTA GAGGTGATCT CTGCTCGAAA GAGAAATAGA ATGAAAATAT

<Whn

-2252 ~~-2302~~ TCTCCGGGCC AGGCGTGGTG GCTCATGCCT GTAATCCCAG CACTTTGGGA

T3R>

SREBP> RAR-α1>

-2202 ~~-2252~~ GGCCAAGGCA TGTGGATCAC CTGAGGTCAG GAGTTCAAAA CCAGCCTGGC

-2152 ~~-2202~~ CAACATGGTG AAACCCCGTC TCTACTAAAA ATACAAAAAA TTAGCTAAGT

-2102 ~~-2152~~ GTGGTGGCGC ATGCCTGTAA TCCCAGCTAC TTGGGAGGGT GAGGCAGGAG

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-1952 AAACAAAAGA ATTAAGCAAA TTAGACATTG CAGAGAGAAC CTGAAGGGGG

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RAR-α1>

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C3

Sac I

-1552 AGCCTGCCTG GTTCTGAGCT CTCATGGTAA GGCTCCTACA GACACGGAAA

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-1452 AGGGCAATAT GGTCCTGAGC AGGGATTAAG AGCTTGGGCT CTCATATGGT

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